

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/889,802

Source: Pct/09

Date Processed by STIC: 8/1/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY.

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 3.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW.

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST-25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

PCT09

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/889,802

DATE: 08/01/2001
TIME: 18:28:48

Input Set : A:\es.txt
Output Set: N:\CRF3\08012001\I889802.raw

pp 1-3
Does Not Comply
Corrected Diskette Needed

w--> 1 **SEQUENZPROTOKOLL** *delete*

3 <110> APPLICANT: Kreutzer Dr., Roland
4 Limmer Dr., Stephan
6 <120> TITLE OF INVENTION: Verfahren und Medikament zur Hemmung der Expression
7 eines vorgegebenen Gens
9 <130> FILE REFERENCE: 400968

c--> 11 <140> CURRENT APPLICATION NUMBER: US/09/889,802

c--> 12 <141> CURRENT FILING DATE: 2001-07-20
14 <150> PRIOR APPLICATION NUMBER: 199 03 713.2
15 <151> PRIOR FILING DATE: 1999-01-30
17 <150> PRIOR APPLICATION NUMBER: 199 56 568.6
18 <151> PRIOR FILING DATE: 1999-11-24

20 <160> NUMBER OF SEQ ID NOS: 8

22 <170> SOFTWARE: PatentIn Ver. 2.1

24 <210> SEQ ID NO: 1

25 <211> LENGTH: 45

26 <212> TYPE: DNA

27 <213> ORGANISM: Knstliche Sequenz

29 <220> FEATURE:

30 <223> OTHER INFORMATION: Beschreibung der knstlichen Sequenz:
31 EcoRI-Schnittstelle, T7-RNA-Polymerasepromotor

33 <400> SEQUENCE: 1

34 ggaattctaa tacgactcac tatagggcga tcagatctct agaag

45

37 <210> SEQ ID NO: 2

38 <211> LENGTH: 50

39 <212> TYPE: DNA

40 <213> ORGANISM: Knstliche Sequenz

42 <220> FEATURE:

43 <223> OTHER INFORMATION: Beschreibung der knstlichen Sequenz:

44 BamHI-Schnittstelle, SP6-RNA-Polymerasepromotor

46 <400> SEQUENCE: 2

47 gggatccatt taggtgacac tatagaatac ccatgatcgc gtagtcgata

50

50 <210> SEQ ID NO: 3

51 <211> LENGTH: 340

52 <212> TYPE: RNA

53 <213> ORGANISM: Knstliche Sequenz

55 <220> FEATURE:

56 <223> OTHER INFORMATION: Beschreibung der knstlichen Sequenz: RNA, die

57 einer Sequenz aus der "positive control DNA" des

58 HeLaScribe Nuclear Extract in vitro

59 Transkriptionskits der Firma Promega entspricht

61 <400> SEQUENCE: 3

62 ucagaucucu agaagcuccua augcgguagu uuaucacagu uaaaauugcua acgcagucag 60

63 gcaccgugua ugaaaucuua caaugcgcuc aucgucaucc uccggcaccgu caccuggau 120

64 gcuguaggca uaggcuuggu uaugccggua cugccgggcc ucuugcgggua uaucguccau 180

65 uccgacagca ucgccaguca cuauggcgug cugcuagcgc uauaugcguu gaugcaauuu 240

66 cuaugcgcac ccguucucgg agcacugucc gaccgcuuug gccgcccggccc aguccugcuc 300

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67 gcuucgcuac uuggagccac uaucgacuac gcgaucaugg 340
 70 <210> SEQ ID NO: 4
 71 <211> LENGTH: 363
 72 <212> TYPE: DNA
 73 <213> ORGANISM: Knstliche Sequenz
 75 <220> FEATURE:
 76 <223> OTHER INFORMATION: Beschreibung der knstlichen Sequenz: DNA, die
 einer Sequenz aus der "positive control DNA" des
 HeLaScribe Nuclear Extract in vitro
 Transkriptionskits der Firma Promega entspricht
 81 <400> SEQUENCE: 4
 82 tcaagatctct agaagttta atgcggtagt ttatcacagt taaattgtcta acgcagtcag 60
 83 gcaccgtgt tgaaatctaa caatgcgtc atcgcatcc tcggcacccgt caccctggat 120
 84 gctgttaggca taggcttgggt tatgccggta ctgcggggcc tcttgcggga tatgtccat 180
 85 tccgacagca tcgcccagtca ctatggcgtg ctgctagcgc tatatgcgtt gatgcaattt 240
 86 ctatgcgcac ccgttctcggt agcactgtcc gaccgctttg gccggccccc agtccctgctc 300
 87 gcttcgctac ttggagccac tatcgactac gcgatcatgg cgaccacacc cgtccctgtgg 360
 88 atc 363
 91 <210> SEQ ID NO: 5
 92 <211> LENGTH: 315
 93 <212> TYPE: RNA
 94 <213> ORGANISM: Knstliche Sequenz
 96 <220> FEATURE:
 97 <223> OTHER INFORMATION: Beschreibung der knstlichen Sequenz: Sequenz aus
 dem YFP-Gen
 100 <400> SEQUENCE: 5
 101 auggugagca agggcgagga gcuguucacc gggguggugc ccauccuggu cgagcuggac 60
 102 ggcgacguaa acggccacaa guucagcgug uccggcgagg gcgaggggcga ugccaccuac 120
 103 ggcaaggcuga cccugaaguu caucugcacc accggcaagc ugccccugcc cuggccccacc 180
 104 cucgugacca cccugaccua cggcgugcag ugcuucagcc gcuaccgcga ccacaugaag 240
 105 cagcacgacu ucuucaaguc cgccaugccc gaaggcuacg uccaggagcg caccacuuc 300
 106 uucaaggacg acggc 315
 109 <210> SEQ ID NO: 6
 110 <211> LENGTH: 52
 111 <212> TYPE: DNA
 112 <213> ORGANISM: Knstliche Sequenz
 114 <220> FEATURE:
 115 <223> OTHER INFORMATION: Beschreibung der knstlichen Sequenz:
 116 EcoRI-Schnittstelle, T7-RNA-Polymerasepromotor,
 117 komplement, rer Bereich zum YFP-Gen
 119 <400> SEQUENCE: 6
 120 ggaattctaa tacgactcac tatagggcga atggtagca agggcgagga gc 52
 123 <210> SEQ ID NO: 7
 124 <211> LENGTH: 53
 125 <212> TYPE: DNA
 126 <213> ORGANISM: Knstliche Sequenz
 128 <220> FEATURE:
 129 <223> OTHER INFORMATION: Beschreibung der knstlichen Sequenz:
 130 BamHI-Schnittstelle, SP6-RNA-Polymerasepromotor,

RAW SEQUENCE LISTING
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131 komplement,er Bereich zum YFP-Gen
133 <400> SEQUENCE: 7
134 gggatccatt tagtgacac tatagaatac gccgtcgcc ttgaagaaga tgg 53
137 <210> SEQ ID NO: 8
138 <211> LENGTH: 21
139 <212> TYPE: RNA
140 <213> ORGANISM Knstliche Sequenz
142 <220> FEATURE:
143 <223> OTHER INFORMATION: Beschreibung der knstlichen Sequenz: RNA, die
144 einer Sequenz aus dem YFP-Gen entspricht
146 <400> SEQUENCE: 8
147 ucgagcugga cggcgacqua a 21

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/889,802

DATE: 08/01/2001

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Input Set : A:\es.txt

Output Set: N:\CRF3\08012001\I889802.raw

L:1 M:259 W: Allowed number of lines exceeded, (1) GENERAL INFORMATION:

L:11 M:270 C: Current Application Number differs, Replaced Application Number

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date

STATISTICS SUMMARY
PATENT APPLICATION: US/09/889,802

DATE: 08/01/2001
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Input Set : A:\es.txt
Output Set: N:\CRF3\08012001\I889802.raw

Application Serial Number: US/09/889,802
Alpha or Numeric: Numeric
Application Class:
Application File Date: 07-20-2001
Art Unit: PCT09
Software Application: PatentIn
Total Number of Sequences: 8
Total Nucleotides: 1239
Total Amino Acids: 0
Number of Errors: 0
Number of Warnings: 1
Number of Corrections: 2

MESSAGE SUMMARY

259 W: 1 (Allowed number of lines exceeded)
270 C: 1 (Current Application Number differs)
271 C: 1 (Current Filing Date differs)